46

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RAW SEQUENCE LISTING PATENT APPLICATION US/08/817,547A

DATE: 04/14/98 TIME: 17:06:41

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This Raw Listing contains the General Information Section and up to the first 5 pages.
 1
 2
 3
            General Information:
     (1)
 4
 5
          (i) APPLICANT:
                             Adermann, Knut
 6
                             Hock, Dieter
 7
                             Magerlein, Markus
 8
 9
         (ii) TITLE OF INVENTION: Peptides from the hPTH Sequence
10
        (iii) NUMBER OF SEQUENCES: 36
11
12
         (iv) CORRESPONDENCE ADDRESS:
13
14
               (A) ADDRESSEE: Jones & Askew, LLP
15
               (B) STREET: 191 Peachtree Street, 37th Floor
16
               (C) CITY: Atlanta
17
               (D) STATE: Georgia
18
               (E) COUNTRY: USA
19
               (F) ZIP: 30303
20
21
          (v) COMPUTER READABLE FORM:
22
                (A) MEDIUM TYPE: Floppy disk
23
                (B) COMPUTER: IBM PC compatible
24
                (C) OPERATING SYSTEM: PC-DOS/MS-DOS
25
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
26
27
         (vi) CURRENT APPLICATION DATA:
                (A) APPLICATION NUMBER: 08/817,547
28
29
               (B) FILING DATE:
30
31
        (vii) PRIOR APPLICATION DATA:
32
                (A) APPLICATION NUMBER: PCT/EP95/03757
33
               (B) FILING DATE: 29 SEPT 1994
34
35
      (viii) ATTORNEY/AGENT INFORMATION:
36
                (A) NAME: FROST, ROGER T.
37
                (B) REGISTRATION NUMBER: 22,176
38
               (C) REFERENCE/DOCKET NUMBER: 07826-0007
39
40
       (ix) TELECOMMUNICATION INFORMATION:
41
                (A) TELEPHONE: 404-818-3700
42
               (B) TELEFAX: 404-818-3799
43
44
45
     (2) INFORMATION FOR SEQ ID NO: 1:
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(i) SEQUENCE CHARACTERISTICS:
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48
               (A) LENGTH: 10 amino acids
49
               (B) TYPE: amino acid
50
               (C) STRANDEDNESS: unknown
51
               (D) TOPOLOGY: unknown
52
53
         (ii) MOLECULE TYPE: peptide
54
55
        (iii) HYPOTHETICAL: no
56
57
        (iv) ANTI-SENSE: no
58
59
60
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
61
62
63
          Ser Val Ser Glu Ile Gln Leu Met His Asn
64
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65
     (2) INFORMATION FOR SEQ ID NO: 2:
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67
     (i) SEQUENCE CHARACTERISTICS:
68
69
               (A) LENGTH: 9 amino acids
70
               (B) TYPE: amino acid
71
               (C) STRANDEDNESS: unknown
72
               (D) TOPOLOGY: unknown
73
74
         (ii) MOLECULE TYPE: peptide
75
76
        (iii) HYPOTHETICAL: no
77
78
        (iv) ANTI-SENSE: no
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80
81
    (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
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83
84
          Ser Val Ser Glu Ile Gln Leu Met His
85
86
87
    (2) INFORMATION FOR SEQ ID NO: 3:
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               (A) LENGTH: 8 amino acids
91
               (B) TYPE: amino acid
92
               (C) STRANDEDNESS: unknown
93
               (D) TOPOLOGY: unknown
94
95
        (ii) MOLECULE TYPE: peptide
96
97
        (iii) HYPOTHETICAL: no
98
99
        (iv) ANTI-SENSE: no
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100
101
102
      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
103
104
105
           Ser Val Ser Glu Ile Gln Leu Met
106
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     (2) INFORMATION FOR SEQ ID NO: 4:
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                (B) TYPE: amino acid
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113
                (C) STRANDEDNESS: unknown
                (D) TOPOLOGY: unknown
114
115
116
         (ii) MOLECULE TYPE: peptide
117
118
         (iii) HYPOTHETICAL: no
119
120
         (iv) ANTI-SENSE: no
121
122
123
124
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
125
126
          Ser Val Ser Glu Ile Gln Leu
127
128
     (2) INFORMATION FOR SEQ ID NO: 5:
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130
131
     (i) SEQUENCE CHARACTERISTICS:
                (A) LENGTH: 6 amino acids
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133
                (B) TYPE: amino acid
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                (C) STRANDEDNESS: unknown
135
                (D) TOPOLOGY: unknown
136
137
         (ii) MOLECULE TYPE: peptide
138
139
         (iii) HYPOTHETICAL: no
140
141
         (iv) ANTI-SENSE: no
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143
144
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
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146
          Ser Val Ser Glu Ile Gln
147
148
149
150
     (2) INFORMATION FOR SEQ ID NO: 6:
151
     (i) SEQUENCE CHARACTERISTICS:
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                (B) TYPE: amino acid
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                (C) STRANDEDNESS: unknown
156
                (D) TOPOLOGY: unknown
157
158
         (ii) MOLECULE TYPE: peptide
159
160
         (iii) HYPOTHETICAL: no
161
162
         (iv) ANTI-SENSE: no
163
164
165
      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
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167
168
           Ser Val Ser Glu Ile
169
           1
                           5
170
     (2) INFORMATION FOR SEQ ID NO: 7:
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173
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175
                (B) TYPE: amino acid
176
                (C) STRANDEDNESS: unknown
177
                (D) TOPOLOGY: unknown
178
         (ii) MOLECULE TYPE: peptide
179
180
181
         (iii) HYPOTHETICAL: no
182
183
         (iv) ANTI-SENSE: no
184
185
186
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
187
188
          His Asn Leu Gly Lys His Leu Asn Ser Met
189
190
                           5
                                                10
191
     (2) INFORMATION FOR SEQ ID NO: 8:
192
193
194
     (i) SEQUENCE CHARACTERISTICS:
195
                (A) LENGTH: 9 amino acids
196
                (B) TYPE: amino acid
                (C) STRANDEDNESS: unknown
197
198
                (D) TOPOLOGY: unknown
199
200
         (ii) MOLECULE TYPE: peptide
201
         (iii) HYPOTHETICAL: no
202
203
204
         (iv) ANTI-SENSE: no
205
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206
207
208
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
209
210
          Asn Leu Gly Lys His Leu Asn Ser Met
211
                           5
212
     (2) INFORMATION FOR SEQ ID NO: 9:
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214
     (i) SEQUENCE CHARACTERISTICS:
215
216
                (A) LENGTH: 8 amino acids
                (B) TYPE: amino acid
217
                (C) STRANDEDNESS: unknown
218
219
                (D) TOPOLOGY: unknown
220
         (ii) MOLECULE TYPE: peptide
221
222
223
        (iii) HYPOTHETICAL: no
224
225
         (iv) ANTI-SENSE: no
226
227
228
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
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230
          Leu Gly Lys His Leu Asn Ser Met
231
232
                           5
233
234
     (2) INFORMATION FOR SEQ ID NO: 10:
235
236
     (i) SEQUENCE CHARACTERISTICS:
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237
                (B) TYPE: amino acid
238
239
                (C) STRANDEDNESS: unknown
240
                (D) TOPOLOGY: unknown
241
242
         (ii) MOLECULE TYPE: peptide
243
        (iii) HYPOTHETICAL: no
244
245
         (iv) ANTI-SENSE: no
246
247
248
249
250
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
251
252
          Gly Lys His Leu Asn Ser Met
253
          1
                           5
254
     (2) INFORMATION FOR SEQ ID NO: 11:
255
256
257
     (i) SEQUENCE CHARACTERISTICS:
258
                (A) LENGTH: 6 amino acids
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SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/817,547A

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Original Text